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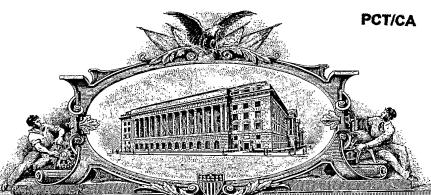
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## PROVISIONAL APPLICATION FOR PATENT COVER SHEET This is a request for filing a PROVISIONAL APPLICATION FOR PATENT under 37 CFR 1.53(c).

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# USE ONLY FOR FILING A PROVISIONAL APPLICATION FOR PATENT

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#### TITLE OF INVENTION

# PREPARATION OF CANOLA PROTEIN ISOLATE AND USE IN AQUACULTURE

#### FIELD OF INVENTION

[0001] The present invention relates to the preparation of canola protein isolates and their use in aquaculture.

#### **BACKGROUND TO THE INVENTION**

Canola protein isolates can be formed from canola oil seed meal. In [0002] copending United States Patent Application No. 10/137,391 filed May 3, 2002 and corresponding PCT Publication No. WO 02/089597, assigned to the assignee hereof and the disclosures of which are incorporated herein by reference, there is described a method of making canola protein isolates from canola oil seed meal, such isolates having at least 100 wt% protein content (N x 6.25). The procedure involves a multiple step process comprising extracting canola oil seed meal using a salt solution, separating the resulting aqueous protein solution from residual oil seed meal, increasing the protein concentration of the aqueous solution to at least about 200 g/L while maintaining the ionic strength substantially constant by using a selective membrane technique, diluting the resulting concentrated protein solution into chilled water to cause the formation of protein micelles, settling the protein micelles to form an amorphous, sticky, gelatinous gluten-like protein micellar mass (PMM), and recovering the protein micellar mass from supernatant having a protein content of at least about 100 wt% as determined by Kjeldahl nitrogen (N x 6.25). As used herein, protein content is determined on a dry weight basis. The recovered PMM may be dried.

[0003] In one embodiment of the process described above and as specifically described in Application No. 10/137,391, the supernatant from the PMM settling step is processed to recover a protein isolate comprising dried protein from the wet PMM and supernatant. This procedure may be effected by initially concentrating the supernatant using ultrafiltration membranes, mixing the concentrated supernatant with the wet PMM and drying the mixture. The resulting canola protein isolate has a high purity of at least about 90 wt% of protein (N x 6.25), preferably at least about 100 wt% protein (N x 6.25).

[0004] In another embodiment of the process described above and as specifically described in Application No. 10/137,391, the supernatant from the PMM settling step is processed to recover a protein from the supernatant. This procedure may be effected by initially concentrating the supernatant using ultrafiltration membranes and drying the concentrate. The resulting canola protein isolate has a high purity of at least about 90 wt% protein (N x 6.25), preferably at least about 100 wt% protein (N x 6.25).

The procedures described in the aforementioned US Patent Applications [0005] are essentially batch procedures. In copending United States Patent Application No. 10/298,678 filed November 19, 2002 and corresponding PCT Publication No. WO 03/043439, assigned to the assignee hereof and the disclosures of which are incorporated herein by reference, there is described a continuous process for making canola protein isolates. In accordance therewith, canola oil seed meal is continuously mixed with a salt solution, the mixture is conveyed through a pipe while extracting protein from the canola oil seed meal to form an aqueous protein solution, the aqueous protein solution is continuously separated from residual canola oil seed meal, the aqueous protein solution is continuously conveyed through a selective membrane operation to increase the protein content of the aqueous protein solution to at least about 200 g/L while maintaining the ionic strength substantially constant, the resulting concentrated protein solution is continuously mixed with chilled water to cause the formation of protein micelles, and the protein micelles are continuously permitted to settle while the supernatant is continuously overflowed until the desired amount of PMM has accumulated in the settling vessel. The PMM is removed from the settling vessel and may be dried. The PMM has a protein content of at least about 90 wt% as determined by Kjeldahl nitrogen (N x 6.25), preferably at least about 100 wt% (N x 6.25).

[0006] As described in the aforementioned US Patent Application No. 10/137,391, the overflowed supernatant may be processed to recover canola protein isolate therefrom.

[0007] Canola seed is known to contain about 10 to about 30 wt% proteins and several different protein components have been identified. These proteins are distinguished by different sedimentation coefficients (S). These known and identified proteins include a 12S globulin, known as cruciferin, and a 2S storage protein, known as napin.

[0008] As described in copending United States Patent Application No. 10/413,371 filed April 15, 2003 and corresponding PCT Publication No. WO 03/088760, assigned to the assignee hereof and the disclosures of which are incorporated herein by references, the PMM-derived canola protein isolate consists predominantly of the 7S protein along with some 12S protein while the supernatant-derived canola protein isolate consists predominantly of the 2S protein.

[0009] In such prior process, canola protein isolates are separately derived from the concentrated canola protein solution by precipitating PMM and separately processing the supernatant to obtain additional quantities of canola protein solution.

[0010] Canola is also known as rapeseed or oil seed rape.

#### SUMMARY OF INVENTION

[0011] In the present invention, the concentrated protein solution resulting from the protein concentration step is dried directly without processing to produce PMM and separately processing the supernatant. This procedure simplifies the production of a canola protein isolate which has a broad spectrum of 12S, 7S and 2S proteins. Because of the lesser number of process steps, the isolate is formed in a more economic manner.

[0012] The canola protein isolate produced according to the process herein may be used in conventional applications of protein isolates, such as, protein fortification of processed foods, emulsification of oils, body formers in baked goods and foaming agents in products which entrap gases. In addition, the canola protein isolates may be formed into protein fibers, useful in meat analogs, may be used as an egg white substitute or extender in food products where egg white is used as a binder. The canola protein isolate may be used as nutritional supplements. Other uses of the canola protein isolate are in pet foods, animal feed, aquaculture and in industrial and cosmetic applications and in personal care products.

[0013] Since the protein isolates which are formed by the process of the present invention are generally of lesser purity, in particular, a higher salt content, than obtained by the procedures described in the aforementioned US patent application, they are preferably used in non-human applications. One particular use of the protein isolates is as a feed in aquaculture, as described in more detail below. However, the protein isolates may be processed to reduce the residual salt content by any convenient procedure, such as by dialysis.

# BRIEF DESCRIPTION OF DRAWINGS

[0014] Figures 1A to 1C are HPLC chromatograms profiles of a canola protein isolate produced in a bench extraction procedure; and

[0015] Figures 2A and 2B are HPLC chromatograms profiles of a canola protein isolate proceeded in a pilot plant scale extraction procedure.

## GENERAL DESCRIPTION OF INVENTION

[0016] The canola protein isolate may be isolated from canola oil seed meal by either a batch process or a continuous process or a semi-continuous process as generally described in the aforementioned United States patent application.

[0017] The initial step of the process of providing the canola protein isolates involves solubilizing proteinaceous material from canola oil seed meal. The proteinaceous material recovered from canola seed meal may be the protein naturally occurring in canola seed or other oil seeds or the proteinaceous material may be a protein modified by genetic manipulation but possessing characteristic hydrophobic and polar properties of the natural protein. The canola meal may be any canola meal resulting from the removal of canola oil from canola oil seed with varying levels of non-denatured protein, resulting, for example, from hot hexane extraction or cold oil extrusion methods. The removal of canola oil from canola oil seed usually is effected as a separate operation from the protein isolate recovery procedure described herein.

[0018] Protein solubilization is effected most efficiently by using a food grade salt solution since the presence of the salt enhances the removal of soluble protein from the oil seed meal. Where the canola protein isolate is intended for non-food uses, non-food-grade chemicals may be used. The salt usually is sodium chloride, although other salts, such as, potassium chloride, may be used. The salt solution has an ionic strength of at least about 0.05, preferably at least about 0.10, to enable solubilization of significant quantities of protein to be effected. As the ionic strength of the salt solution increases, the degree of solubilization of protein in the oil seed meal initially increases until a maximum value is achieved. Any subsequent increase in ionic strength does not increase the total protein solubilized. The ionic strength of the food grade salt solution which causes maximum protein solubilization varies depending on the salt concerned and the oil seed meal chosen.

[0019] In view of the greater degree of dilution required for protein precipitation with increasing ionic strengths, it is usually preferred to utilize an ionic strength value less than about 0.8, and more preferably a value of about 0.1 to about 0.6.

[0020] In a batch process, the salt solubilization of the protein is effected at a temperature of at least about 5°C and preferably up to about 35°C, preferably accompanied by agitation to decrease the solubilization time, which is usually about 10 to about 60 minutes. It is preferred to effect the solubilization to extract substantially as much protein from the oil seed meal as is practicable, so as to provide an overall high product yield.

[0021] The lower temperature limit of about 5°C is chosen since solubilization is impractically slow below this temperature while the upper preferred temperature limit of about 35°C is chosen since the process becomes uneconomic at higher temperature levels in a batch mode.

[0022] In a continuous process, the extraction of the protein from the canola oil seed meal is carried out in any manner consistent with effecting a continuous extraction of protein from the canola oil seed meal. In one embodiment, the canola oil seed meal is continuously mixed with a food grade salt solution and the mixture is conveyed through a pipe or conduit having a length and at a flow rate for a residence time sufficient to effect the desired extraction in accordance with the parameters described herein. In such continuous procedure, the salt solubilization step is effected rapidly, in a time of up to about 10 minutes, preferably to effect solubilization to extract substantially as much protein from the canola oil seed meal as is practicable. The solubilization in the continuous procedure preferably is effect at elevated temperatures, preferably above about 35°C, generally up to about 65°C.

[0023] The aqueous food grade salt solution and the canola oil seed meal have a natural pH of about 5 to about 6.8 to enable a protein isolate to be formed by the micellar route, as described in more detail below.

[0024] At and close to the limits of the pH range, protein isolate formation occurs only partly through the micelle route and in lower yields than attainable elsewhere in the pH range. For these reasons, pH values of about 5.3 to about 6.2 are preferred.

[0025] The pH of the salt solution may be adjusted to any desired value within the range of about 5 to about 6.8 for use in the extraction step by the use of any convenient acid, usually hydrochloric acid, or alkali, usually sodium hydroxide, as required.

[0026] The concentration of oil seed meal in the food grade salt solution during the solubilization step may vary widely. Typical concentration values are about 5 to about 15% w/v.

[0027] The protein extraction step with the aqueous salt solution has the additional effect of solubilizing fats which may be present in the canola meal, which then results in the fats being present in the aqueous phase.

[0028] The protein solution resulting from the extraction step generally has a protein concentration of about 5 to about 40 g/L, preferably about 10 to about 30 g/L.

[0029] The aqueous phase resulting from the extraction step then may be separated from the residual canola meal, in any convenient manner, such as by employing vacuum filtration, followed by centrifugation and/or filtration to remove residual meal. The separated residual meal may be dried for disposal.

[0030] The colour of the final canola protein isolate can be improved in terms of light colour and less intense yellow by the mixing of powdered activated carbon or other pigment adsorbing agent with the separated aqueous protein solution and subsequently removing the adsorbent, conveniently by filtration, to provide a protein solution. Diafiltration also may be used for pigment removal.

[0031] Such pigment removal step may be carried out under any convenient conditions, generally at the ambient temperature of the separated aqueous protein solution, employing any suitable pigment adsorbing agent. For powdered activated carbon, an amount of about 0.025% to about 5% w/v, preferably about 0.05% to about 2% w/v, is employed.

[0032] Where the canola seed meal contains significant quantities of fat, as described in US Patents Nos. 5,844,086 and 6,005,076, assigned to the assignee hereof and the disclosures of which are incorporated herein by reference, then the defatting steps described therein may be effected on the separated aqueous protein solution and on the concentrated aqueous protein solution discussed below. When the colour improvement step is carried out, such step may be effected after the first defatting step.

[0033] As an alternative to extracting the oil seed meal with an aqueous salt solution, such extraction may be made using water alone, although the utilization of water alone tends to extract less protein from the oil seed meal than the aqueous salt solution. Where such alternative is employed, then the salt, in the concentrations discussed above, may be added to the protein solution after separation from the residual oil seed meal in order to maintain the protein in solution during the concentration step described below. When a colour removal step and/or a first fat removal step is carried out, the salt generally is added after completion of such operations.

[0034] Another alternative procedure is to extract the oil seed meal with the food grade salt solution at a relatively high pH value above about 6.8, generally up to about 9.9. The pH of the food grade salt solution, may be adjusted in pH to the desired alkaline value by the use of any convenient food-grade alkali, such as aqueous sodium hydroxide solution. Alternatively, the oil seed meal may be extracted with the salt solution at a relatively low pH below about pH 5, generally down to about pH 3. Where such alternative is employed, the aqueous phase resulting from the oil seed meal extraction step then is separated from the residual canola meal, in any convenient manner, such as by employing vacuum filtration, followed by centrifugation and/or filtration to remove residual meal. The separated residual meal may be dried for disposal.

[0035] The aqueous protein solution resulting from the high or low pH extraction step then is pH adjusted to the range of about 5 to about 6.8, preferably about 5.3 to about 6.2, as discussed above, prior to further processing as discussed below. Such pH adjustment may be effected using any convenient acid, such as hydrochloric acid, or alkali, such as sodium hydroxide, as appropriate.

[0036] The aqueous protein solution then is concentrated, usually about 4 to about 20 fold, to increase the protein concentration thereof while maintaining the ionic strength thereof substantially constant. Such concentration generally is effected to provide a concentrated protein solution having a protein concentration of at least about 50 g/L, preferably at least about 200 g/L, more preferably at least about 250 g/L.

[0037] The concentration step may be effected in any convenient manner consistent with batch or continuous operation, such as by employing any convenient selective membrane technique, such as ultrafiltration or diafiltration, using membranes, such as hollow-fibre membranes or spiral-wound membranes, with a suitable molecular

weight cut-off, such as about 3000 to about 50,000 daltons, preferably about 5000 to about 10,000 daltons, having regard to differing membrane materials and configurations. The membranes may be hollow-fibre or spiral-wound. For continuous operation, the membranes may be dimensioned to permit the desired degree of concentration as the aqueous protein solution passes through the membranes.

[0038] The concentrated protein solution then may be subjected to a diafiltration step using an aqueous salt solution of the same molarity and pH as the extraction solution. Such diafiltration may be effected using from about 2 to about 20 volumes of diafiltration solution, preferably about 5 to about 10 volumes of diafiltration solution. In the diafiltration operation, further quantities of contamination are removed from the aqueous protein solution by passage through the membrane with the permeate. The diafiltration operation may be effected until no significant further quantities of phenolics and visible colour are present in the permeate. Such diafiltration may be effected using a membrane having a molecular weight cut-off in the range of about 3000 to about 50,000 daltons, preferably about 5,000 to about 10,000 daltons, having regard to different membrane materials and configuration.

[0039] An antioxidant may be present in the diafiltration medium using at least part of the diafiltration step. The antioxidant may be any convenient food grade antioxidant, such as sodium sulfite or ascorbic acid. The quantity of antioxidant employed in the diafiltration medium depends on the materials employed and may vary from about 0.01 to about 1 wt%, preferably about 0.05 wt%. The antioxidant serves to inhibit oxidation of phenolics present in the concentrated canola protein isolate solution.

[0040] The concentration step and the diafiltration step may be effected at any convenient temperature, generally about 20° to about 60°C, and for the period of time to effect the desired degree of concentration. The temperature and other conditions used to some degree depend upon the membrane equipment used to effect the concentration and the desired protein concentration of the solution.

[0041] The concentrating of the protein solution to the preferred concentration above about 200 g/L in this step not only increases the process yield to levels above about 40% in terms of the proportion of extracted protein which is recovered as dried protein isolate, preferably above about 80%, but also decreases the salt concentration of the final protein isolate after drying. The ability to control the salt concentration of the

isolate is important in applications of the isolate where variations in salt concentrations affect the functional and sensory properties in a specific food application.

[0042] As is well known, ultrafiltration and similar selective membrane techniques permit low molecular weight species to pass therethrough while preventing higher molecular weight species from so doing. The low molecular weight species include not only the ionic species of the food grade salt but also low molecular weight materials extracted from the source material, such as, carbohydrates, pigments and antinutritional factors, as well as any low molecular weight forms of the protein. The molecular weight cut-off of the membrane is usually chosen to ensure retention of a significant proportion of the protein in the solution, while permitting contaminants to pass through having regard to the different membrane materials and configurations.

[0043] The concentrated and optionally diafiltered protein solution may be subject to a further defatting operation, if required, as described in US Patents Nos. 5,844,086 and 6,005,076.

[0044] The concentrated and optionally diafiltered protein solution may be subject to a colour removal operation as an alternative to the colour removal operation described above. Powdered activated carbon may be used herein as well as granulated activated carbon (GAC). Another material which may be used as a colour adsorbing agent is polyvinyl pyrrolidone.

The colour absorbing agent treatment step may be carried out under any convenient conditions, generally at the ambient temperature of the canola protein solution. For powdered activated carbon, an amount of about 0.025% to about 5% w/v, preferably about 0.05% to about 2% w/v, may be used. Where polyvinylpyrrolidone is used as the colour adsorbing agent, an amount of about 0.5 to about 5 w/v, preferably about 2 to about 3% w/v, may be used. The colour adsorbing agent may be removed from the canola protein solution by any convenient means, such as by filtration.

from the option colour removal step may be subjected to pasteurization to kill any bacteria which may have been present in the original meal as a result of storage or otherwise and extracted from the meal into the canola protein isolate solution in the extraction step. Such pasteurization may be effected under any desired pasteurization conditions. Generally, the concentrated and optionally diafiltered protein solution is

heated to a temperature of about 55° to about 70°C, preferably about 60° to about 65°C, for about 10 to about 15 minutes, preferably about 10 minutes. The pasteurized concentrated protein solution then may be cooled for further processing as described below, preferably to a temperature of about 25° to about 40°C.

[0047] The concentrated protein solution resulting from the concentration step, optional diafiltration step, optional colour removal step and optional defatting step then is dried by any convenient technique, such as spray drying, freeze drying or vacuum drum drying, to a dry form to provide a canola protein isolate having a protein content of at least about 90 wt% protein  $(N \times 6.25)$  and is substantially undenatured, (as determined by differential scanning calorimetry).

[0048] As mentioned previously, one potential use of the canola protein isolate is in aquaculture. In Mwachireya et al, Aquaculture Nutrition 5(2), 73-82, it was observed that a canola protein product produced by the PMM process provided growth rates, feed efficiencies and protein efficiency ratio for rainbow trout and Atlantic salmon superior to prior art canola protein materials.

[0049] In terms of protein digestibility, the canola protein product had a protein digestibility coefficient better than fish meal, an energy digestibility coefficient similar to fish meal and a calculated digestible energy similar to fish meal. The canola protein product, even at lower than optimum protein concentrations, showed the same growth ratio and feed intake as a commercial feed material.

[0050] The protein efficiency ratio (PER) is the single most important positive indicator for all the protein preparations. The canola protein product used in the studies reported in Mwachireya et al had less than optimal protein concentrations resulting from processing difficulties but, nevertheless the canola protein isolate (CPI-L) diet had a comparable PER to a basal diet which was a special research diet and a special commercial diet (RCC). The PER of the RCC diet is statistically the same as the CPI-L and basal PER values.

[0051] Having regard to the protein distribution in the product of the invention and the provision of a true protein isolate, it is expected that improved feeding results, compared to those achieved by Mwachireya et al, can be achieved for Salmonids.

[0052] When the canola protein isolate is formed by drying of the concentrated protein solution, the product contains significantly greater concentration of residual salt

than isolation via the PMM procedure discussed in the aforementioned prior art. The presence of the salt is not detrimental to certain uses of the protein isolate, for example, in the use in aquaculture.

[0053] However, where the presence of the salt is detrimental to the intended use of the canola protein isolate, salt may be removed by dialyzing an aqueous solution of the protein, which may be in the form of the concentrated, optionally diafiltered, prior to drying.

#### **EXAMPLES**

#### Example 1:

[0054] This Example illustrates the procedure of the invention for the provision of canola protein isolates.

[0055] 150 kg of commercial canola oil seed meal lot AL022 was added to 1010.5L 0.1M saline (NaCl) at 19.8°C and mixed for 30 minutes to provide an aqueous protein solution. At the halfway point of mixing (15 minutes), 0.05 wt% or 500g w/v of ascorbic acid was added as an antioxidant. Extraction pH was 6.12 with no adjustment being made to the natural pH of the saline.

[0056] In order to remove the meal from the extracted solution, meal slurry was passed over a vacuum filter belt and a solution of 790L with an average protein content of 1.74 wt% (17.4 g/L) was the result.

[0057] This solution was then passed through a desludger centrifuge and filter press housing 2.0 um pads in order to further clarify the protein solution. The final clarified protein extract had a volume of 780L and a protein content of 1.58 wt% (15.8 g/L).

[0058] A 700L aliquot of the clarified protein solution was then ultrafiltered on a 2-membrane system using polyvinyldiene difluoride (PVDF) 5 spiral wound membranes. These membranes have a MWCO range of 5000 Daltons. Total volume reduction was from 700L down to 32L or 21.8 times volume reduction. The resulting 32L of concentrated protein solution or retentate had an average protein content of 25.10 wt% (251 g/L).

[0059] The retentate from the UF step was pasteurized at 60°C for 10 minutes and aliquots were then dried on an APV spray dryer.

[0060] Final protein content of the dried product was 93.08 wt% as is and on dry weight basis 95.46 wt% (N x 6.25). Percentage nitrogen values were determined using a Leco FP528 Nitrogen Determinator). The batch was designated BW-AL022-I02-03A.

#### Example 2:

[0061] This Example describes the preparation of a laboratory scale sample of canola protein isolate.

[0062] 75 g of the same canola meal as used in Example 1 was added to 500 mL of 0.10 M saline solution (15% w/w) and the mixture was shaken for 30 minutes at 220 rpm on a rotational shaker. The extract, containing 1.99 wt% protein, was centrifuged for 20 minutes at 10,000 rpm and filtered through crepe-fluted filter paper.

[0063] 350 ml of filtrate was concentrated on an Amicon Ultrafiltration unit using a 5,000 MWCO PES membrane until 150 ml of retentate was collected. The retentate was diafiltered with 0.1 M saline solution to produce 75 ml of DF retentate containing 6.24 wt% protein.

[0064] The retentate was dialyzed using Spectra/Por 6 to 8,000 MWCO tubing at refrigerated temperature. The dialyzed sample was frozen and then freeze-dried. The resulting canola protein isolate had a protein content of 101 wt% (N x 6.25).

#### Example 3:

[0065] This Example provides protein analysis of the canola protein isolates produced in Examples 1 and 2.

[0066] HPLC analysis was conducted on the canola protein isolates prepared as described in Examples 1 and 2. HPLC chromatograms of the bench extract, bench DF permeate and the bench DF UF dialyzed canola protein isolate are shown in Figures 1A, 1B and 1C, respectively.

[0067] HPLC chromatograms of the BW-AL022-I02-03A samples on two different dates are shown in Figures 2A and 2B.

Analysis of the canola protein isolates prepared as described in Examples 1 and 2 is contained in Tables 1 to 5 below. Table 5 contains amino acid analysis of the samples in comparison to typical PMM-derived (C200) and supernatant derived (C300) canola protein isolates, prepared as described in copending US patent application No. 10/266,701 filed October 9, 2002, assigned to the assignee hereof and the disclosure of which is incorporated herein by reference.

[0069] As may be seen from this data, the bench isolate (Example 2) shows a higher protein ratio than the I02 isolate (Example 1), based on peak areas. Both indicate that the globular proteins (7S, 12S, >12S, and sub-unit) comprise about 2/3rd of the total protein peak areas, with the albumins (2S and pronapin) contributing the other 1/3rd.

[0070] Other components found in the HPLC chromatograms indicate a relatively higher phytate level in the bench isolate, with a lower phenolic (and miscellaneous) content, based on peak areas. This result indicated that the bench isolate contained less free phenolic acid content than the I02 isolate. Colour differences, based on A<sub>330</sub> may be due to bound phenolics on the protein, which are not removable on the filter membranes.

[0071] The I02 HPLC-SEC (Figure 2, Table 2) profile remained largely unchanged from the initial scan made on September 19<sup>th</sup>, 2003, to the more current run on December 18<sup>th</sup>, 2003, with the exception of ascorbic acid. Ascorbic acid oxidizes over time and was reduced in quantity, as determined by peak area, over this time frame. As a result, the other components increased in ratio, as shown, but this had little effect on the protein ratios.

[0072] The HPLC-SEC analyses of the bench samples, (Extract, UF Permeate, DF Permeate, DF Retentate and DF dialyzed FD Retentate resolubilized) (Figure 1, Table 1), indicates that UF, DF and dialysis steps removed the majority of the phenolics and miscellaneous components, but was less effective with phytic acid removal. Phytic acid tends to have a strong association with protein. Even so, phytic acid was observed in the bench permeate HPLC chromatograms, which indicates partial removal through the membranes, (perhaps 20 to 30% of the total).

[0073] The Example 1 isolate contained salt and other minerals as shown, amounting to about 3% of the final dry weight of isolate (Table 3). No toxic elements were detected. The results show that the bench isolate is higher in protein, due to the DF and dialysis steps used in the preparation of this sample.

[0074] The amino acid analysis results were converted to "grams per 100 grams amino acids" in Tables 4A and 4B. The averages and standard deviations are also shown and indicate minimal differences. This is expected since the DF and dialysis steps remove non-proteins, which would not affect the amino acid balance in any significant way unless there were a lot of free amino acids and peptides.

[0075] Table 5 compares the current sample amino acid profiles with results from earlier studies. The current retentates are very similar in composition to the earlier retentates (from A8 and A10 meals) as well as a Puratein sample from A10 meal. Puratein is a mixture of PMM and Supertein and should be similar to the retentate analysis.

[0076] Table 5 also shows C200 and C300 (A10 meal) amino acid profiles. The retentate and Puratein samples fall between these two isolates, which would be expected.

[0077] Lysine is an essential amino acid that is in low abundance for cereals. Oilseeds, particularly canola, tend to have higher levels of lysine. The retentate analysis reveals a significant amount of lysine and this would improve the nutritional quality of this isolate, (even for fish or other non-human feed). The essential acid composition is quite high for retentate isolate, as is shown at the bottom of Table 5.

[0078] Overall, analysis shows that retentate, C500, is an isolate with a quality amino acid composition that ranges between C200 and C300. This isolate has low levels of non-proteins, and the bench study shows that salts, phenolics and other unknown substances are removed by Ultra-filtration. Diafiltration and dialysis can improve this elimination of non-proteins, as shown by the bench extraction data. However, this is not required to produce an isolate, as is shown by the I02 results.

#### SUMMARY OF DISCLOSURE

[0079] In summary of the disclosure, there is provided a novel process for the preparation of oil seed isolates having multiple uses, including in aquaculture. Modifications are possible within the scope of the invention.

#### Table 1

Burcon NutraScience Winnipeg Lab C500 Sample for External Analysis December 19, 2003

RP30 27 C500 Dec19 03.xls

Lab UF DF FD C500 Lab # 21,681
Extraction started on Dec. 11 by Brandy Gosnell and FD retentate removed on Wednesday, Dec. 17/03.

Moisture: Leco protein (as is): Leco protein (dry basis):	17-Dec Average: n.d. 101.09% n.d.	s.d.: n.d. 0.03% n.d.		Terms:  MWCO = mo cutoff  n.d. = not don FD = freeze-c DF = diafilter PES = polyeti	ie Iried ed		
HPLC-SEC:  Protein % of Total Area: Phytate % of Total Area: Phenolics % of Total Area: Misc. % of Total Area: 12S+>12S	Diluted 1:4 21,670 Extraction 23.8% 3.3% 37.6% 35.3% 6.5%	neat 21,679 Permeate 0.0% 5.2% 52.2% 42.6% 0.0%	neat 21,680 DF Perm. 0.0% 4.3% 47.6% 48.1% 0.0%	Diluted 1:4 21,681 DF retent. 62.6% 11.2% 12.7% 13.5% 6.9%	1% W/V sol'n 21,681 DF FD ret. 77.3% 8.7% 3.6% 10.4% 6.7%	of protein area	
7S. sub-unit 2S+ pronapin globulins % of total:	58.0% 0.2% 35.4% 64.6%	0.0% 0.0% 0.0% 0.0%	0.0% 0.0% 0.0% 0.0%	57.2% 0.1% 35.9% 64.1%	57.7% 0.2% 35.5% 64.5%	11 11 11	11 13 21 17

#### Miscellaneous:

A330 3.70 AU on 1% W/V sample in water = 80.7 ug/mL Sinapine equivalent

= 0.81% on dry sample weight

pH = 5.83

Sent to POS, Saskatoon on Dec. 17th for amino acid analysis.

#### Table 2

#### BURCON NUTRASCIENCE WINNIPEG LAB C500 SAMPLES FOR EXTERNAL ANALYSIS

December 19, 2003

AL022 102 C500 Analysis Dec19 03.xls

BW-AL022-I02-03A #1 C500	Lab # 20,576												
submitted for analysis on Sept. 18/0	3 and revised Nov I	8/03	*										
Analyses Date:	18-Sep		Dec										
Allaryses Date.	Average:	s.d.:		rage:	s.d.:								
Moisture:	2.49%	0.06%		.d.	. n.d.								
Leco protein (as is):	93.08%	0.24%		57%	0.50%								
Leco protein (dry basis):	95.46%	0.24%	n	ı.d.	n.d.								
Ecco protein (all) easily.	n.e	d. = not done											
HPLC-SEC:		•											
•	1% W/V Sampl	e in 0.1M Na	Cl:										
HPLC Test done on:	19-Sep	18-Dec											
Protein % of Total Area:	72.5%	73.0%											
Phytate % of Total Area:	3.2%	5.0%											
Ascorbate % of Total Area:	4.6%	0.7%	loss o	ver time									
Phenolics % of Total Area:	16.2%	17.3%											
Misc. % of Total Area:	3.5%	3.9%											
12S+>12S	5.5%	5.3%		tein area									
7S	64.1%	62.7%	. "	**									
75 2S	24.0%	24.8%	**	**									
sub-unit	0.1%	1.4%	11	**									
	6.3%	5.9%	tı	**									
pronapin globulins % of total:	69.7%	69.3%	**	"									
globums 70 of total	es of ascorbic acid:												

#### Miscellaneous:

A330 1.45 AU on 1% W/V sample in water
= 31.6 ug/mL Sinapine equivalent
= 0.32% on dry sample weight

Solutes gained in peak area with loss of ascorbic acid.

Sent to POS, Saskatoon on Dec. 17th for amino acid analysis.
Sent to Central Testing on Dec. 17th for moisture, protein and elemental analysis. See Sheet 2.

Table 3

Central Testing Laboratories Report Dated Dec. 23/03

BW-AL022-102-03A #1 C500 Test Results

Lab # 20,576

•		•	
	Sample as rec'd:	Sample Dry Basis	
Moisture Dry Matter Crude Protein (N x 6.25)	3.93% 93.00%	96.07% 96.81%	
Calcium Phosphorus Magnesium Potassium Copper Sodium	0.10% 0.40% 0.11% 0.31% 0.0011% 0.73%	0.10% 0.42% 0.11% 0.32% 0.0011% 0.76%	
Sodium Chloride equivalent Zinc Manganese Iron Boron	1.85% 0.0007% <0.0001% 0.0119% 0.0005%	1.92% 0.0007% <0.0001% 0.0124% 0.0005%	
Lead Cadmium . Mineral sum (1)	0.00 0.00 2.78%	0.00 0.00 · 2.92%	mg/kg mg/kg

<sup>(1)</sup> includes estimate of chloride for sodium. Red denotes toxic elements which are below the threshold limits.

Table 4A

		. Sec	THE STANDARD CARE THE TANK THE STANDARD	0.0	0.04	0.12	0.0	1.13	0.23	0.19	0.31	81.0	5.0°	0.26	0.50	0:03	0.27	0.01	0.49	0.23	0.19	on dry weight basis (DWB)	
			Average	7.25	2.85	3.40	7.33	20.90	4.76	3.89	2.25	5.03	1.71	3.86	7.02	2.09	3.82	7.08	4.32	6.61	6.70	89.87	03:05
ummary	g/100 g dry matter	Dec 15/03 Bench	一十二十二十二十二十二十二十二十二十二十二十二十二十二十二十二十二十二十二十二	7.46	2.82	3.48	1.38	21.70	4.92	4.02	2.47	5.15	1.84	4.04	7.37	2.07	4.01	2.08	4.67	6.77	83	200	93.00
ntate Amino Acid St		3	#20,576	7.04	2.87	3.31	1.28	20.10	4.59	3.75	2.03	4.90	1.57	3.67	99.9	2.11	3,63	202	207			0.30	86.56
Burson NutraScience Ganola Retentate Amino Acid Summary	Analysis: POS Results: January 5/04	Amino	Acid	Aspartic	Threonine	Serine	Tryptophan	Glutamic	Glycine	Alanine	Cystine	Valine	Methionine	Isoleucine	leucine	Tvrosine	Dhenvdalanine	Liction	orie: I	Lysine	Arginine	Proline	Sum:
Rureon NutraS	Analysis POS	Amino Acid	NW:	1	110.1	105.1	204.2	146.1	75.1	89.1	121.1	117.1	149.2	134.0	131.5	4. 6.	101.2.	7:001	155.2	146.2	174.2	115.1	

Note that the two samples are protein isolates, based on crude protein (N  $\times$  6.25) and not on amino acid analyses. Amino acid analyses usually results in loss of some nitrogen through deamination of glutamine and asparagine.

		A			•	Previou	Previous amino acid tests:	d.tests:	
	Amino Acid Summa	nmary: g/100g Amilino Acids	202	SE CANTOCCACAGA SE	S. C. Par	Diratein	Retentate	Retentate	
Amino Acid	Amino	BW-AL022102-03A#1	Dec. 15/03 Bench		Day.	LTA10	A8	A10.04	
MM	Acid	1#20,576	#21/08/1	WALE GOT	2 - 2	7.0	76	7.1	Aspartic *
133.1	Aspartic*		8.0	ö	- :	2 6	· ·	~~~	Thronnine
7 0 7	Throoning	23	3.0	3.2	0.2	5.0	0.0	9 (	
	allicolline.		17	80	0.1	3.9	3.9	4.0	Serine
105.1	Serine	0 . 1	, L		0	1.4	1.2	1.5	Tryptophan <sup>e</sup>
204.2	Tryptophan "	1.5	c.T		3 6	22.5	22.8	20.9	Glutamic *
146.1	Glutamic *	23.2	23.3	6.03		5.4	5.2	5.3	Glycine
75.1	Glycine	5,3	5.3			. Y	4.5	. 97	Alanine
89.1	Alanine	4.3	4.3	4.3	) )	? 1	2 6	00	Overline <sup>8</sup>
1211	Cystine	2.3	2.7	2.5	0.7	7.7	777	ייי ני ו	V-line
	1/-ti== 0	7.7	rc.	5.6	0.1	9.2	2.7		Valine
177.1	Valine		2 6	9	0.1	2.1	6:	6:	Methionine
149.2	Methionine	×2.	קר	<u>;</u> ;		44	4.5	4.5	Soleucine
131.2	Isoleucine	4.2	5.4		- 6	10	7.0	0 8	1 eucine
131.2	Leucine	7.7	7.9	7.8	0.7	0.7	} ;	, ,	Tvrosine
	Tvrosine	2.4	2.2	2.3	0.2	7.3	<b>7.</b>	3 9	1).com
7101	. y.com.c	73	43	4.3	0.1	4.1	4.2	4.2	Phenylalanine
165.2	Phenylalanine			22	0.7	3.2	2.7	333	Histidine "
155.2	Histidine	2,4	7.7	3 19		"	6.7	5.5	Lysine
146.2	Lysine	4,6	5.0	<b>9</b>	. ·	. t	- 77	7.2	Arcinine <sup>0</sup>
174.2	Arginine	7.5	. 7.3	7.4	- G	: :		- V Z	Proline
145.1	Proline	9.2	7.3	7.5	0.2	2,	2 8	*00*	
	Sum:	100.0	100.0	100.0		1007	0.25	40.	
	Sum essential aa:	_	45.8	45.5		47.5	4.0.4	20.0	

e = 11 essential amino acids aa = amino acids
• Glutamic acid and aspartic acid are mostly deaminated glutamine and asparagine.

ae Summary Retentates Jan05 04.xls

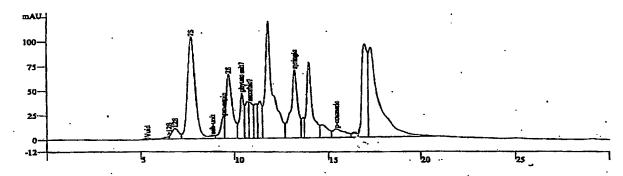
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	C200 TO C300	A10	5.4 10.0	3.7 4.1	3.0		1.4	22.8 19.0	4.9 5.6	4.7 4.7	3.4 1.2		5.4	2.1 1.6	4.2 5.0	-	0.0	2.0 2.8	3.8 4.9	3.6 2.6	3.6	0.0	6.7 7.8	8.2 6.7	100.1 100.1	
	Refentate		_		9 9	0.4	1.5	20.9	5.3	4.6	000	6:7	5.7	1.9		}.	8.0	2.3	4.2	. "	ر ا آ	5.5	7.2	7.4	1007	
	Dotomtato	Aetellidie	7 6	9 6	0°5	ა. მ	1.2	22.8	5.2	, i r	? 6	7.7	5.7	7.0	: '	C.4.	2.9	2.4	-	, t	7.7	4.9	7.4	: ^	8 60	22.0
		Puratein	TO TO	0.7	8. 8.	3.9	14	20.5	7.7	- 14	t (	2.7	5.6		7.7	4.4	7.8			- <del>,</del>	3.2	ຕ	. ^	: :	5.7	1,001
			il.	 	3.2	38		C. C	5.5.5	? .	5.4	2.5	ŭ	9 (		4.3	10	. c	<b>7.7</b>	4.3	2.3	48	) \ \	4. 1	6.7	100:00
	o Acids	Dec. 15/03 Bench	#	0.	3.0	2.7	- I	1,5	23.3	5.3	4.3	2.7		c.	2.0	4.3		S: /	2.2	4.3	2.2	2	n'c	7.3	7.3	100.00
•	ry: g/100g Amin	AL022-102-03A #1	#20,576	8.1	63		5.0 0	3.	23.2	5.3	4.3		) ! !	2.7	1.8		1 r	7.7	2.4	4.2	P C	r (	0.	7.5	7.6	100.00
	Amino Acid Summary: g/100g Amino Acids	Amino	Acid	Aspartic *	0 001110	lucomie	Serine	Tryptophan	Glutamic *	Glycine	Alanine	9	Cystine	Valine	Methionine <sup>9</sup>	8	Isoleucine	Leucine <sup>®</sup>	Tyrosine	Phenylalanine <sup>e</sup>	9 17-17-11	Histiaine	Lysine	Arginine	Proline	Sum:

Current analyses slightly on low side for threonine and histidine and slightly higher for glutamic acid. Overall, analyses are very similar and lie between typical analyses for C200 and C300, as expected. Blue denotes historical analyses of samples done for canola extractions. Puratein is blend of C200 and C300, close to the expected composition of C500.

#### FIGURE 1: Bench Study HPLC-SEC Chromatograms

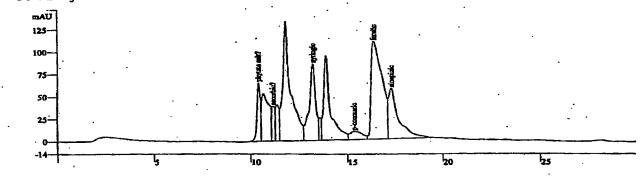
Bench Extract December 11, 2003: SEC-HPLC chromatogram 1:4 dil. 50-uL injection.



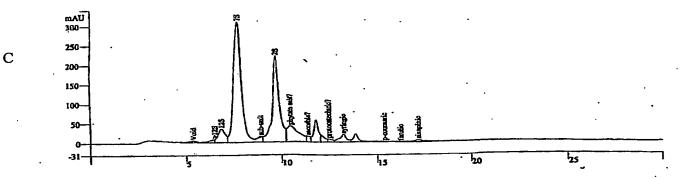
Bench DF Permeate December 12, 2003: SEC-HPLC chromatogram no dil. 50-uL injection.

Α

**B** .

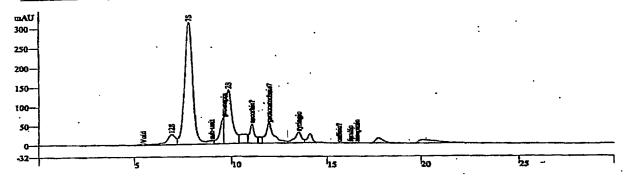


Bench DF UF Dialyzed Isolate December 17, 2003: SEC-HPLC chromatogram 1% 50-uL injection.

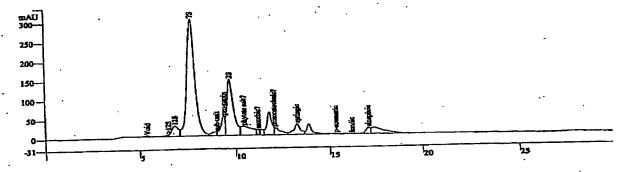


# FIGURE 2: BW-AL022-I02-03A #1 C500 HPLC-SEC Chromatograms

102 Isolate September 19, 2003: SEC-HPLC chromatogram 1% 50-uL injection



102 Isolate December 18, 2003: SEC-HPLC chromatogram 1% 50-uL injection



All samples prepared with 0.10M NaCl.

Α

 $\mathbf{B}$ 

# INITIAL INFORMATION DATA SHEET

#### **Inventor Information:**

Inventor One Given Name:

To follow

Family Name:

Postal Address Line One:

City:

State or Province: Postal or Zip Code: Citizenship Country:

Inventor Two Given Name:

To follow

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City:

State or Province: Postal or Zip Code: Citizenship Country:

# **Correspondence Information**

Correspondence Customer Number: 24,223

## **Application Information**

Title Line One: Title Line Two:

PREPARATION OF CANOLA PROTEIN ISOLATE

AND USE IN AQUACULTURE

Total Drawing Sheets:

Two (2)

Application Type:

Utility Patent

Docket Number:

7865-101 MIS:jb

#### Representative Information

Registration Number:

24,973

#### **Continuity Information**

This application is a: Application One:

Filing Date: